

#9

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/511,814
Source: PCR
Date Processed by STIC: 12/14/05

ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 12/14/2005

PATENT APPLICATION: US/10/511,814

TIME: 14:45:13

Input Set : A:\211080016U2.TXT

Output Set: N:\CRF4\12142005\J511814.raw

4 <110> APPLICANT: McCance, Dennis
 5 Westbrook, III, Thomas F.
 7 <120> TITLE OF INVENTION: E7 REGULATION OF P21 (CIP1) THROUGH AKT
 10 <130> FILE REFERENCE: 21108.0016U2
 12 <140> CURRENT APPLICATION NUMBER: 10/511,814
 C--> 13 <141> **CURRENT FILING DATE: 2004-10-19**
 15 <150> PRIOR APPLICATION NUMBER: PCT/US03/12667
 16 <151> PRIOR FILING DATE: 2003-04-21
 18 <150> PRIOR APPLICATION NUMBER: 60/374,245
 19 <151> PRIOR FILING DATE: 2002-04-19
 21 <160> NUMBER OF SEQ ID NOS: 21
 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 25 <210> SEQ ID NO: 1
 26 <211> LENGTH: 273
 27 <212> TYPE: PRT
 28 <213> ORGANISM: Artificial Sequence
 30 <220> FEATURE:
 31 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
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 34 <400> SEQUENCE: 1
 35 Ser Thr Thr Ser Pro Thr Glu Glu Thr Thr Gln Lys Leu Thr Val Ser
 36 1 5 10 15
 37 His Ile Glu Gly Tyr Glu Cys Gln Pro Ile Phe Leu Asn Val Leu Glu
 38 20 25 30
 39 Ala Ile Glu Pro Gly Val Val Cys Ala Gly His Asp Asn Asn Gln Pro
 40 35 40 45
 41 Asp Ser Phe Ala Ala Leu Leu Ser Ser Leu Asn Glu Leu Gly Glu Arg
 42 50 55 60
 43 Gln Leu Val His Val Val Lys Trp Ala Lys Ala Leu Pro Gly Phe Arg
 44 65 70 75 80
 45 Asn Leu His Val Asp Asp Gln Met Ala Val Ile Gln Tyr Ser Trp Met
 46 85 90 95
 47 Gly Leu Met Val Phe Ala Met Gly Trp Arg Ser Phe Thr Asn Val Asn
 48 100 105 110
 49 Ser Arg Met Leu Tyr Phe Ala Pro Asp Leu Val Phe Asn Glu Tyr Arg
 50 115 120 125
 51 Met His Lys Ser Arg Met Tyr Ser Gln Cys Val Arg Met Arg His Leu
 52 130 135 140
 53 Ser Gln Glu Phe Gly Trp Leu Gln Ile Thr Pro Gln Glu Phe Leu Cys
 54 145 150 155 160
 55 Met Lys Ala Leu Leu Leu Phe Ser Ile Ile Pro Val Asp Gly Leu Lys
 56 165 170 175
 57 Asn Gln Lys Phe Phe Asp Glu Leu Arg Met Asn Tyr Ile Lys Glu Leu

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58          180          185          190
59 Asp Arg Ile Ile Ala Cys Lys Arg Lys Asn Pro Thr Ser Cys Ser Arg
60          195          200          205
61 Arg Phe Tyr Gln Leu Thr Lys Leu Leu Asp Ser Val Gln Pro Ile Ala
62          210          215          220
63 Arg Glu Leu His Gln Phe Thr Phe Asp Leu Leu Ile Lys Ser His Met
64 225          230          235          240
65 Val Ser Val Asp Phe Pro Glu Met Met Ala Glu Ile Ile Ser Val Gln
66          245          250          255
67 Val Pro Lys Ile Leu Ser Gly Lys Val Lys Pro Ile Tyr Phe His Thr
68          260          265          270
69 Gln
72 <210> SEQ ID NO: 2
73 <211> LENGTH: 344
74 <212> TYPE: PRT
75 <213> ORGANISM: Artificial Sequence
77 <220> FEATURE:
78 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
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81 <400> SEQUENCE: 2
82 Trp Ser Gln Pro Lys Thr Pro Val Pro Ala Gln Arg Glu Arg Ala Pro
83 1          5          10          15
84 Val Ser Gly Thr Gln Glu Lys Asn Lys Ile Arg Pro Arg Gly Gln Arg
85          20          25          30
86 Asp Ser Ser Tyr Tyr Trp Glu Ile Glu Ala Ser Glu Val Met Leu Ser
87          35          40          45
88 Thr Arg Ile Gly Ser Gly Ser Phe Gly Thr Val Tyr Lys Gly Lys Trp
89          50          55          60
90 His Gly Asp Val Ala Val Lys Ile Leu Lys Val Val Asp Pro Thr Pro
91 65          70          75          80
92 Glu Gln Phe Gln Ala Phe Arg Asn Glu Val Ala Val Leu Arg Lys Thr
93          85          90          95
94 Arg His Val Asn Ile Leu Leu Phe Met Gly Tyr Met Thr Lys Asp Asn
95          100          105          110
96 Leu Ala Ile Val Thr Gln Trp Cys Glu Gly Ser Ser Leu Tyr Lys His
97          115          120          125
98 Leu His Val Gln Glu Thr Lys Phe Gln Met Phe Gln Leu Ile Asp Ile
99          130          135          140
100 Ala Arg Gln Thr Ala Gln Gly Met Asp Tyr Leu His Ala Lys Asn Ile
101 145          150          155          160
102 Ile His Arg Asp Met Lys Ser Asn Asn Ile Phe Leu His Glu Gly Leu
103          165          170          175
104 Thr Val Lys Ile Gly Asp Phe Gly Leu Ala Thr Val Lys Ser Arg Trp
105          180          185          190
106 Ser Gly Ser Gln Gln Val Glu Gln Pro Thr Gly Ser Val Leu Trp Met
107          195          200          205
108 Ala Pro Glu Val Ile Arg Met Gln Asp Asn Asn Pro Phe Ser Phe Gln
109          210          215          220
110 Ser Asp Val Tyr Ser Tyr Gly Ile Val Leu Tyr Glu Leu Met Thr Gly

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111 225          230          235          240
112 Glu Leu Pro Tyr Ser His Ile Asn Asn Arg Asp Gln Ile Ile Phe Met
113          245          250          255
114 Val Gly Arg Gly Tyr Ala Ser Pro Asp Leu Ser Lys Leu Tyr Lys Asn
115          260          265          270
116 Cys Pro Lys Ala Met Lys Arg Leu Val Ala Asp Cys Val Lys Lys Val
117          275          280          285
118 Lys Glu Glu Arg Pro Leu Phe Pro Gln Ile Leu Ser Ser Ile Glu Leu
119          290          295          300
120 Leu Gln His Ser Leu Pro Lys Ile Asn Arg Ser Ala Ser Glu Pro Ser
121 305          310          315          320
122 Leu His Arg Ala Ala His Thr Glu Asp Ile Asn Ala Cys Thr Leu Thr
123          325          330          335
124 Thr Ser Pro Arg Leu Pro Val Phe
125          340
127 <210> SEQ ID NO: 3
128 <211> LENGTH: 20
129 <212> TYPE: PRT
130 <213> ORGANISM: Artificial Sequence
132 <220> FEATURE:
133 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
134     Synthetic Construct
136 <400> SEQUENCE: 3
137 Lys Met Ser Lys Asp Gly Lys Lys Lys Lys Lys Lys Thr Lys Thr Lys
138 1          5          10          15
139 Cys Ile Ile Met
140          20
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143 <211> LENGTH: 164
144 <212> TYPE: PRT
145 <213> ORGANISM: Artificial Sequence
147 <220> FEATURE:
148 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
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151 <400> SEQUENCE: 4
152 Met Ser Glu Pro Ala Gly Asp Val Arg Gln Asn Pro Cys Gly Ser Lys
153 1          5          10          15
154 Ala Cys Arg Arg Leu Phe Gly Pro Val Asp Ser Glu Gln Leu Arg Arg
155          20          25          30
156 Asp Cys Asp Ala Leu Met Ala Gly Cys Ile Gln Glu Ala Arg Glu Arg
157          35          40          45
158 Trp Asn Phe Asp Phe Val Thr Glu Thr Pro Leu Glu Gly Asp Phe Ala
159          50          55          60
160 Trp Glu Arg Val Arg Gly Leu Gly Leu Pro Lys Leu Tyr Leu Pro Thr
161 65          70          75          80
162 Gly Pro Arg Arg Gly Arg Asp Glu Leu Gly Gly Gly Arg Arg Pro Gly
163          85          90          95
164 Thr Ser Pro Ala Leu Leu Gln Gly Thr Ala Glu Glu Asp His Val Asp
165          100          105          110

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166 Leu Ser Leu Ser Cys Thr Leu Val Pro Arg Ser Gly Glu Gln Ala Glu
167      115      120      125
168 Gly Ser Pro Gly Gly Pro Gly Asp Ser Gln Gly Arg Lys Arg Arg Gln
169      130      135      140
170 Thr Ser Met Thr Asp Phe Tyr His Ser Lys Arg Arg Leu Ile Phe Ser
171 145      150      155      160
172 Lys Arg Lys Pro
175 <210> SEQ ID NO: 5
176 <211> LENGTH: 495
177 <212> TYPE: PRT
178 <213> ORGANISM: Artificial Sequence
180 <220> FEATURE:
181 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
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184 <400> SEQUENCE: 5
185 Ala Thr Gly Thr Cys Ala Gly Ala Ala Cys Cys Gly Gly Cys Thr Gly
186 1      5      10      15
187 Gly Gly Gly Ala Thr Gly Thr Cys Cys Gly Thr Cys Ala Gly Ala Ala
188      20      25      30
189 Cys Cys Cys Ala Thr Gly Cys Gly Cys Cys Ala Gly Cys Ala Ala Gly
190      35      40      45
191 Gly Cys Cys Thr Gly Cys Cys Gly Cys Cys Gly Cys Cys Thr Cys Thr
192      50      55      60
193 Thr Cys Gly Gly Cys Cys Cys Ala Gly Thr Gly Gly Ala Cys Ala Gly
194 65      70      75      80
195 Cys Gly Ala Gly Cys Ala Gly Cys Thr Gly Ala Gly Cys Cys Gly Cys
196      85      90      95
197 Gly Ala Cys Thr Gly Thr Gly Ala Thr Gly Cys Gly Cys Thr Ala Ala
198      100      105      110
199 Thr Gly Gly Cys Gly Gly Gly Cys Thr Gly Cys Ala Thr Cys Cys Ala
200      115      120      125
201 Gly Gly Ala Gly Gly Cys Cys Cys Gly Thr Gly Ala Gly Cys Gly Ala
202      130      135      140
203 Thr Gly Gly Ala Ala Cys Thr Thr Cys Gly Ala Cys Thr Thr Thr Gly
204 145      150      155      160
205 Thr Cys Ala Cys Cys Gly Ala Gly Ala Cys Ala Cys Cys Ala Cys Thr
206      165      170      175
207 Gly Gly Ala Gly Gly Gly Thr Gly Ala Cys Thr Thr Cys Gly Cys Cys
208      180      185      190
209 Thr Gly Gly Gly Ala Gly Cys Gly Thr Gly Thr Gly Cys Gly Gly Gly
210      195      200      205
211 Gly Cys Cys Thr Thr Gly Gly Cys Cys Thr Gly Cys Cys Cys Ala Ala
212      210      215      220
213 Gly Cys Thr Cys Thr Ala Cys Cys Thr Thr Cys Cys Cys Ala Cys Gly
214 225      230      235      240
215 Gly Gly Gly Cys Cys Cys Cys Gly Gly Cys Gly Ala Gly Gly Cys Cys
216      245      250      255
217 Gly Gly Gly Ala Thr Gly Ala Gly Thr Thr Gly Gly Gly Ala Gly Gly
218      260      265      270

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```

219 Ala Gly Gly Cys Ala Gly Gly Cys Gly Gly Cys Cys Thr Gly Gly Cys
220      275      280      285
221 Ala Cys Cys Thr Cys Ala Cys Cys Thr Gly Cys Thr Cys Thr Gly Cys
222      290      295      300
223 Thr Gly Cys Ala Gly Gly Gly Gly Ala Cys Ala Gly Cys Ala Gly Ala
224 305      310      315      320
225 Gly Gly Ala Ala Gly Ala Cys Cys Ala Thr Gly Thr Gly Gly Ala Cys
226      325      330      335
227 Cys Thr Gly Thr Cys Ala Cys Thr Gly Thr Cys Thr Thr Gly Thr Ala
228      340      345      350
229 Cys Cys Cys Thr Thr Gly Thr Gly Cys Cys Thr Cys Gly Cys Thr Cys
230      355      360      365
231 Ala Gly Gly Gly Gly Ala Gly Cys Ala Gly Gly Cys Thr Gly Ala Ala
232      370      375      380
233 Gly Gly Gly Thr Cys Cys Cys Cys Ala Gly Gly Thr Gly Gly Ala Cys
234 385      390      395      400
235 Cys Thr Gly Gly Ala Gly Ala Cys Thr Cys Thr Cys Ala Gly Gly Gly
236      405      410      415
237 Thr Cys Gly Ala Ala Ala Cys Gly Gly Cys Gly Gly Cys Ala Gly
238      420      425      430
239 Ala Cys Cys Ala Gly Cys Ala Thr Gly Ala Cys Ala Gly Ala Thr Thr
240      435      440      445
241 Thr Cys Thr Ala Cys Cys Ala Cys Thr Cys Cys Ala Ala Ala Cys Gly
242      450      455      460
243 Cys Cys Gly Gly Cys Thr Gly Ala Thr Cys Thr Thr Cys Thr Cys Cys
244 465      470      475      480
245 Ala Ala Gly Ala Gly Gly Ala Ala Gly Cys Cys Cys Thr Ala Ala
246      485      490      495
248 <210> SEQ ID NO: 6
249 <211> LENGTH: 480
250 <212> TYPE: PRT
251 <213> ORGANISM: Artificial Sequence
253 <220> FEATURE:
254 <223> OTHER INFORMATION: Description of Artificial Sequence:/Note =
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260 Glu Tyr Ile Lys Thr Trp Arg Pro Arg Tyr Phe Leu Leu Lys Asn Asp
261      20      25      30
262 Gly Thr Phe Ile Gly Tyr Lys Glu Arg Pro Gln Asp Val Asp Gln Arg
263      35      40      45
264 Glu Ala Pro Leu Asn Asn Phe Ser Val Ala Gln Cys Gln Leu Met Lys
265      50      55      60
266 Thr Glu Arg Pro Arg Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp
267 65      70      75      80
268 Thr Thr Val Ile Glu Arg Thr Phe His Val Glu Thr Pro Glu Glu Arg
269      85      90      95
270 Glu Glu Trp Thr Thr Ala Ile Gln Thr Val Ala Asp Gly Leu Lys Lys

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date